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FIGURE 1

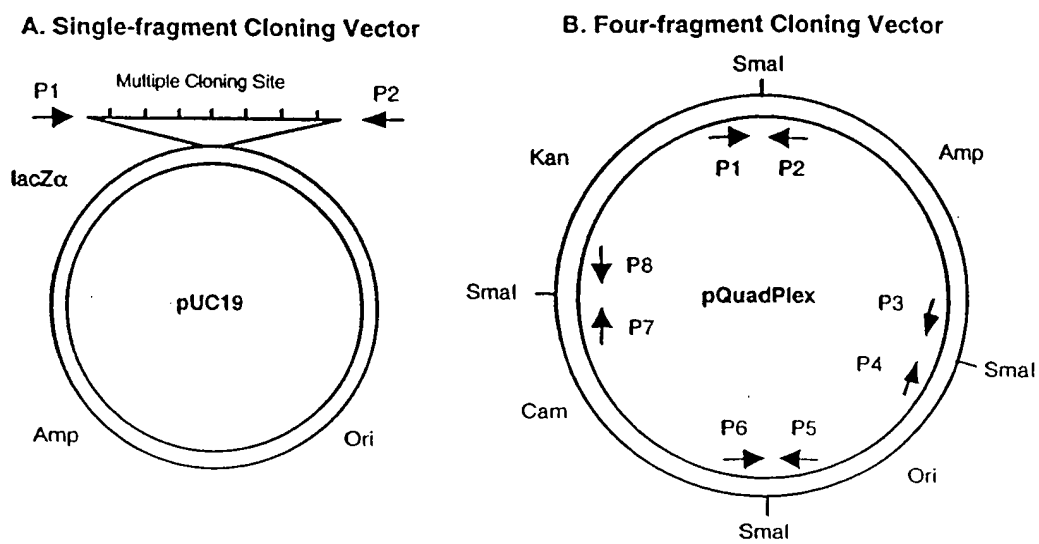


FIGURE 2

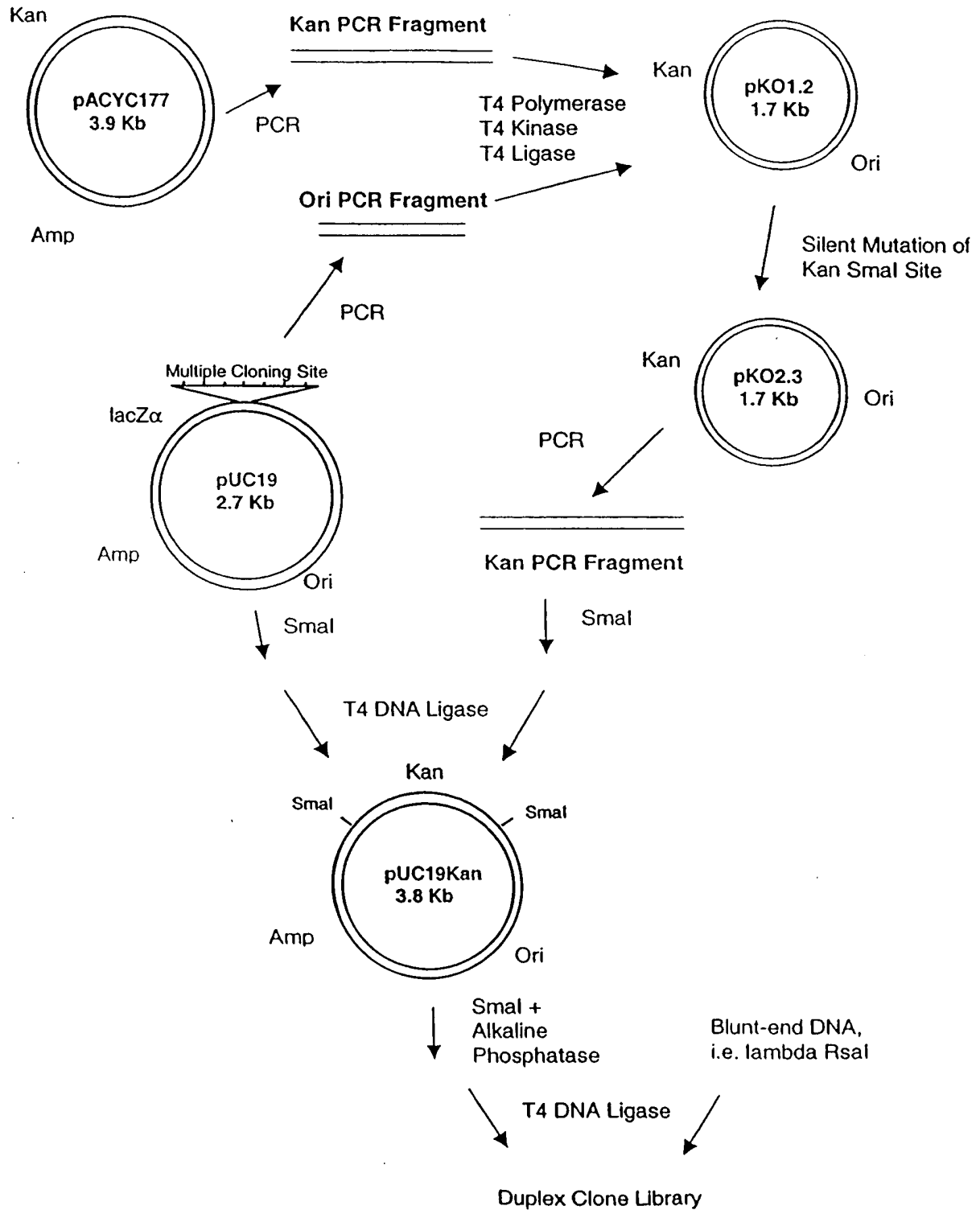


FIGURE 3

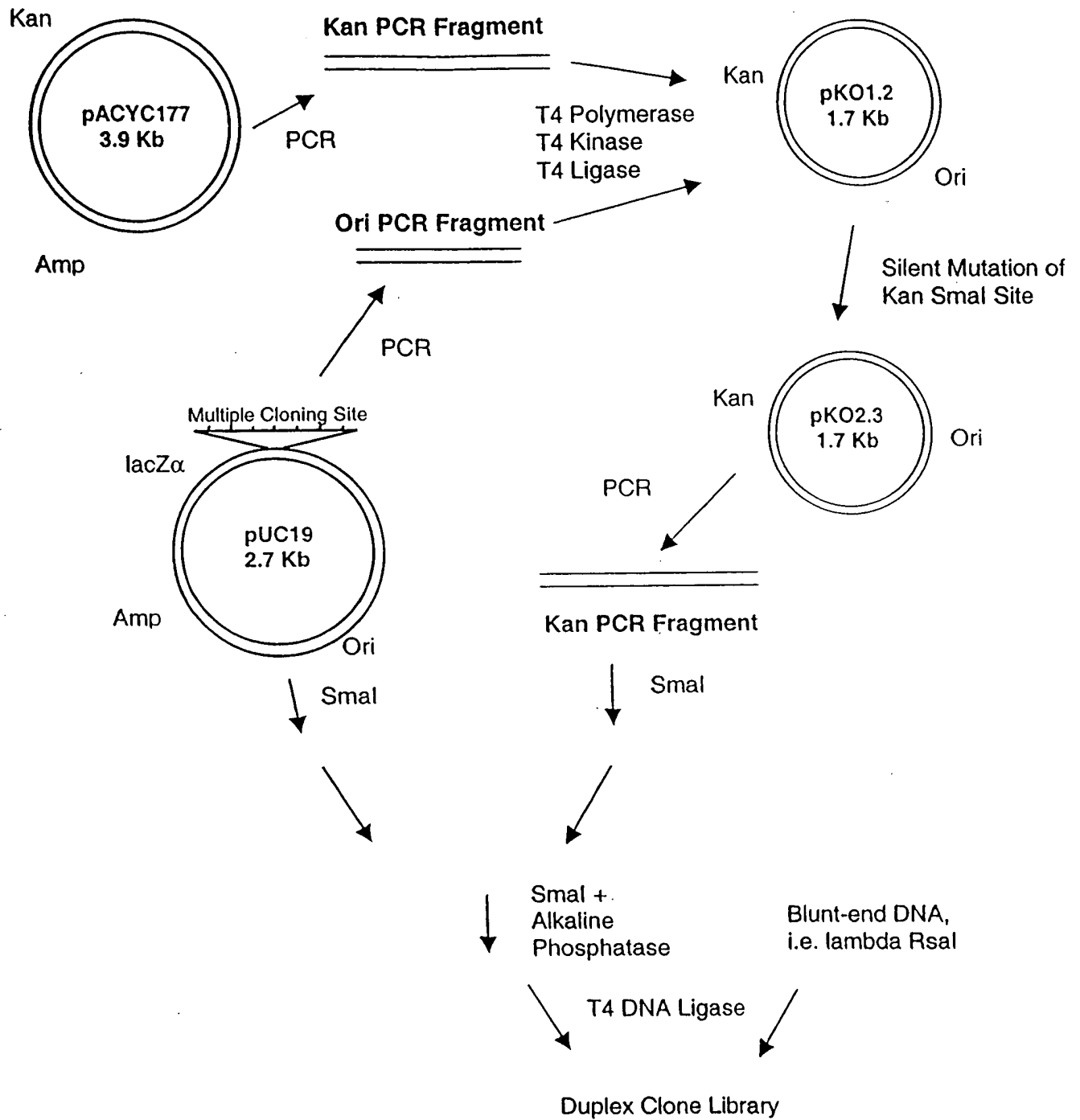


FIGURE 4

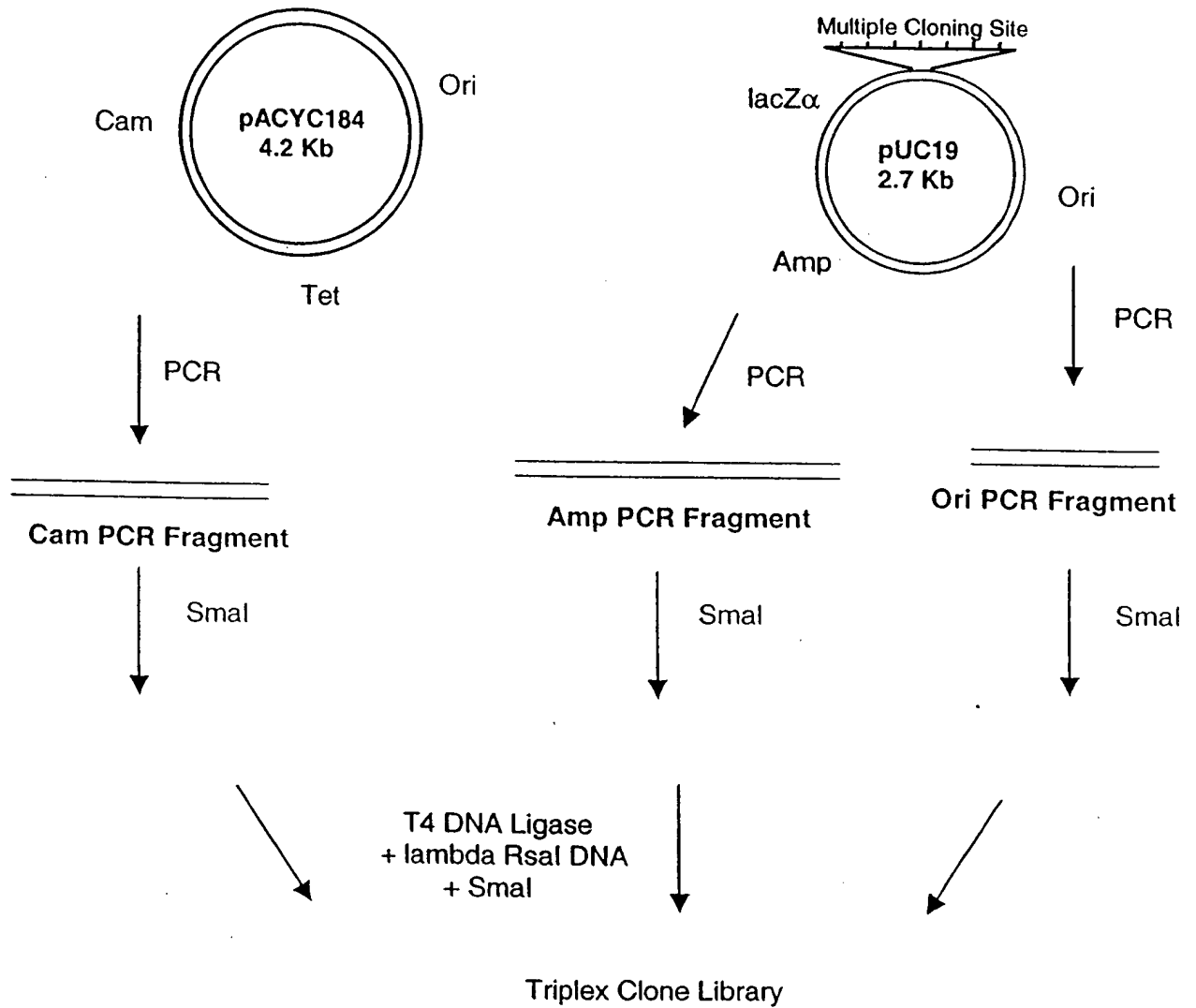


FIGURE 5

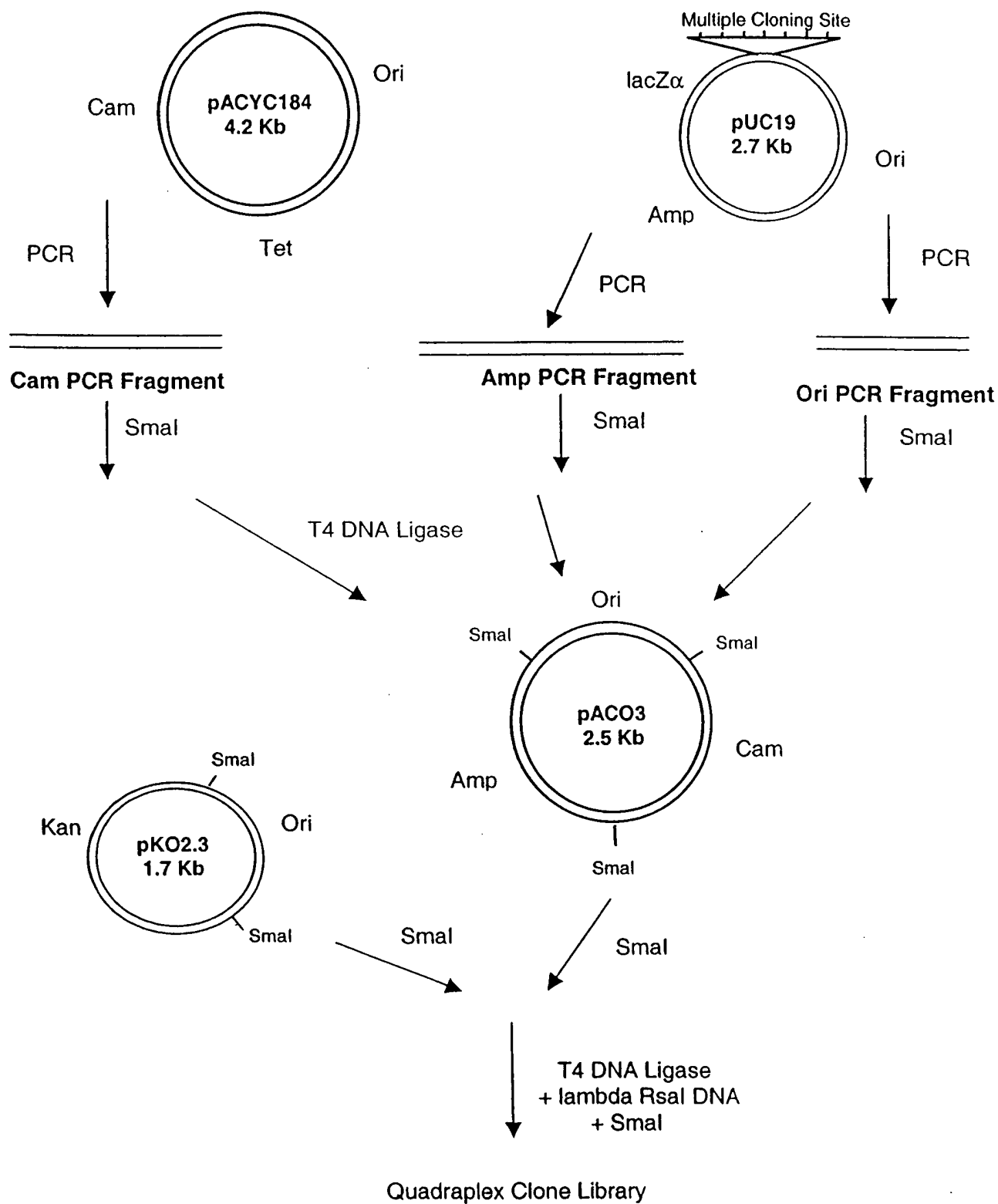


FIGURE 6

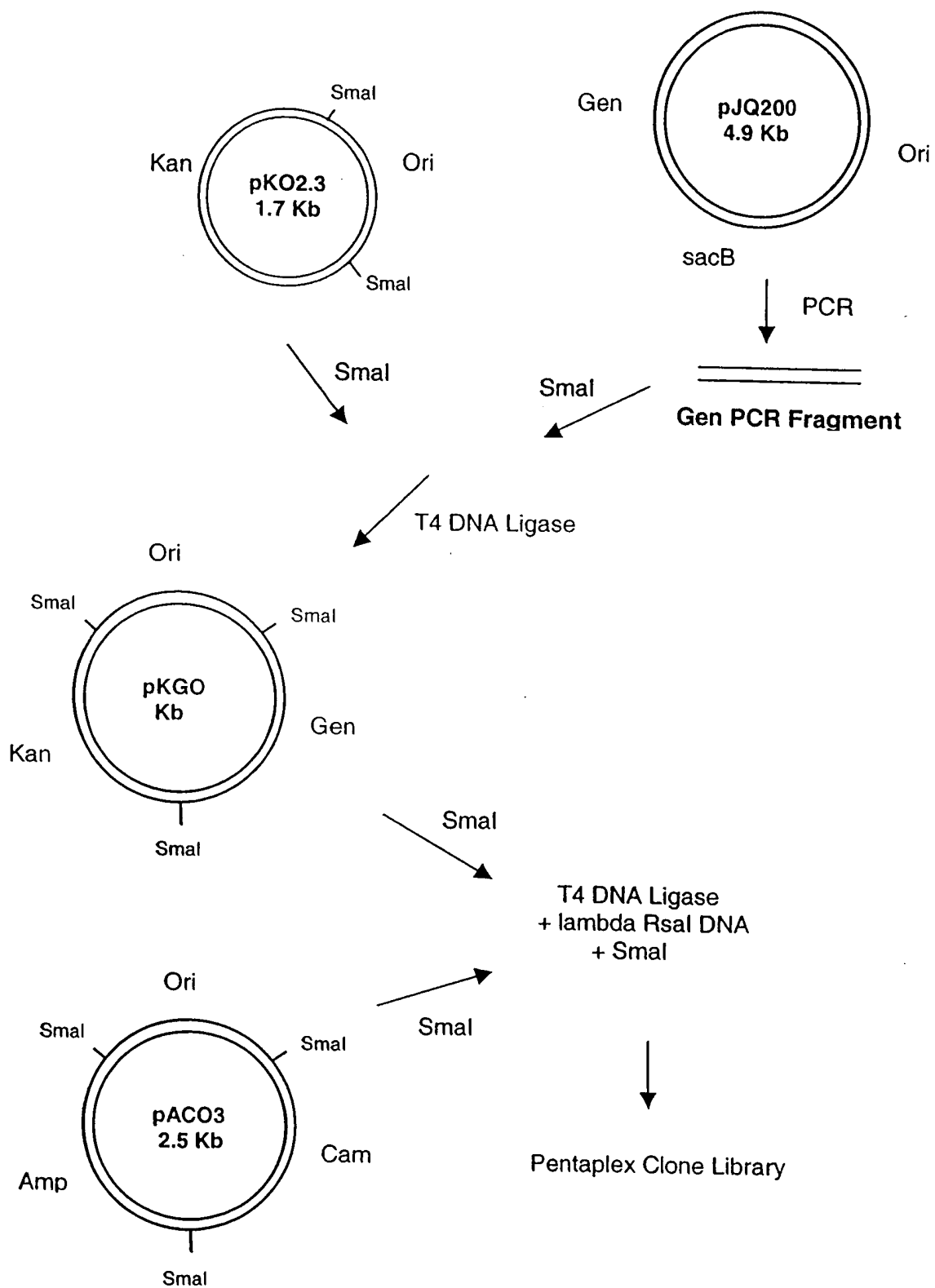


FIGURE 7

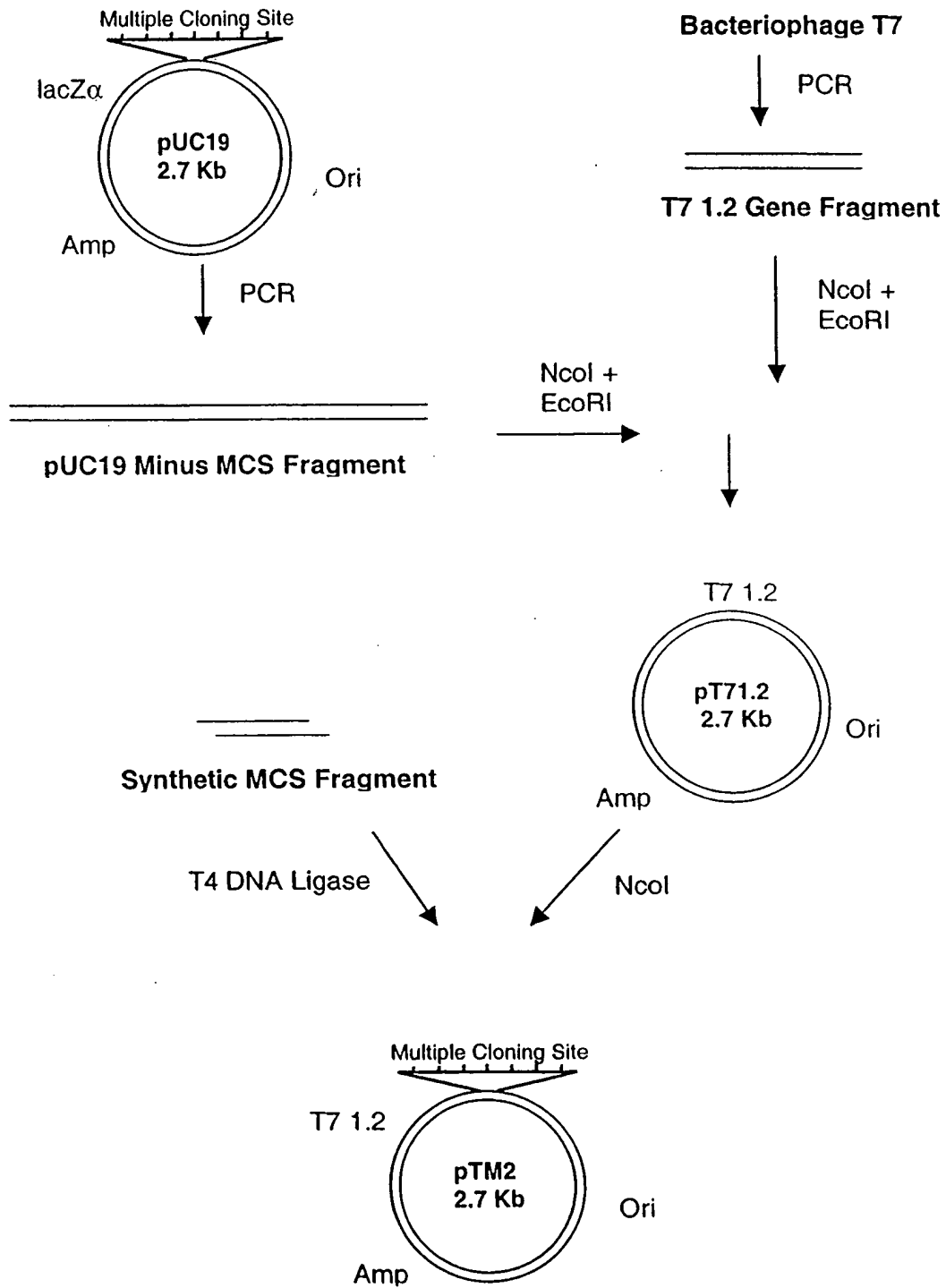


FIGURE 8

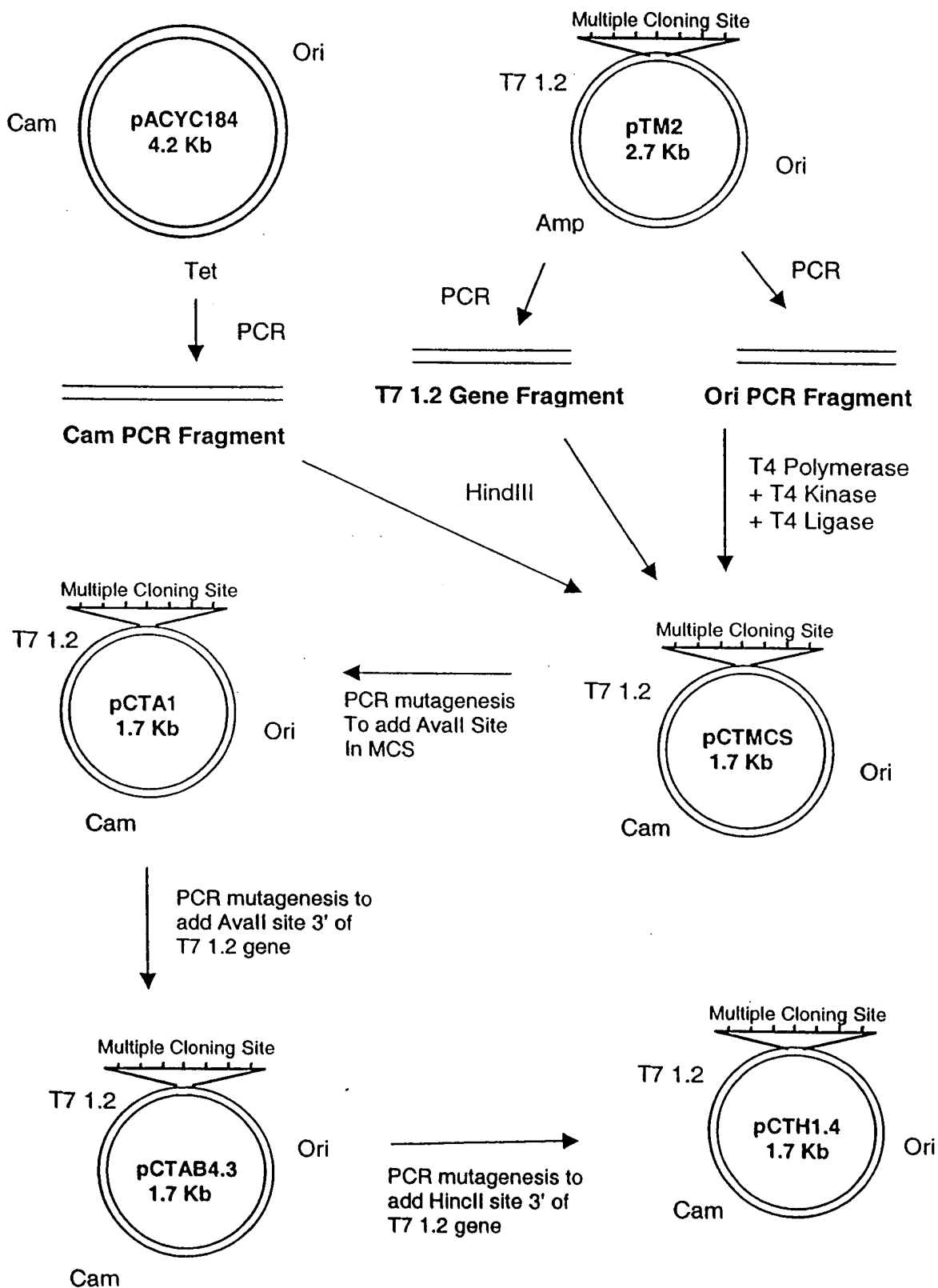


FIGURE 9

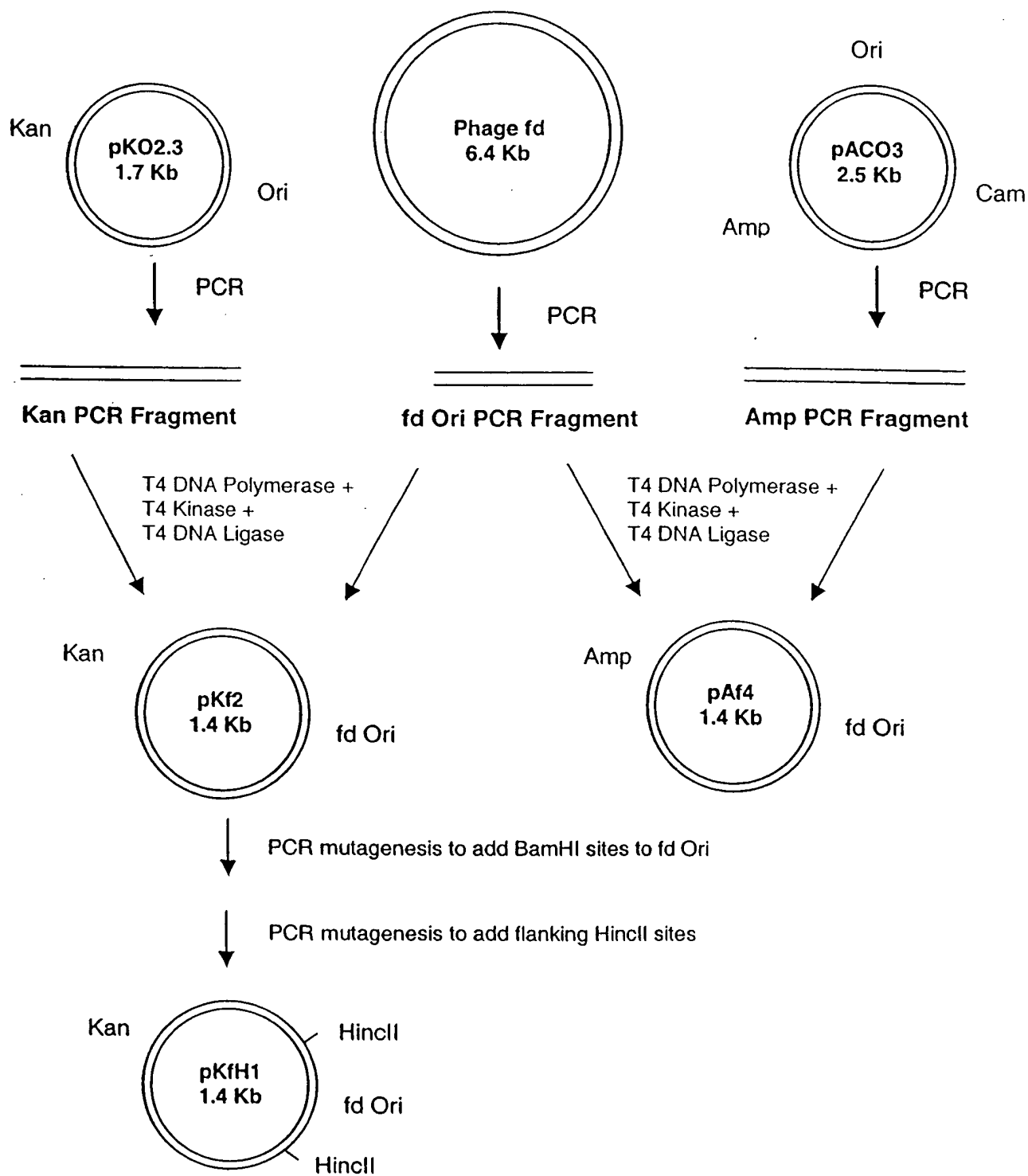


FIGURE 10

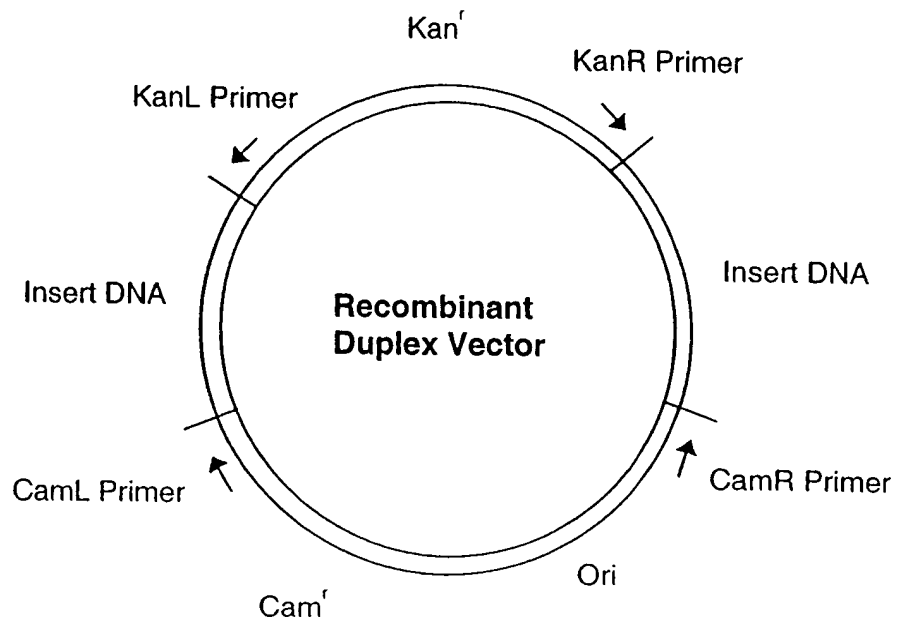
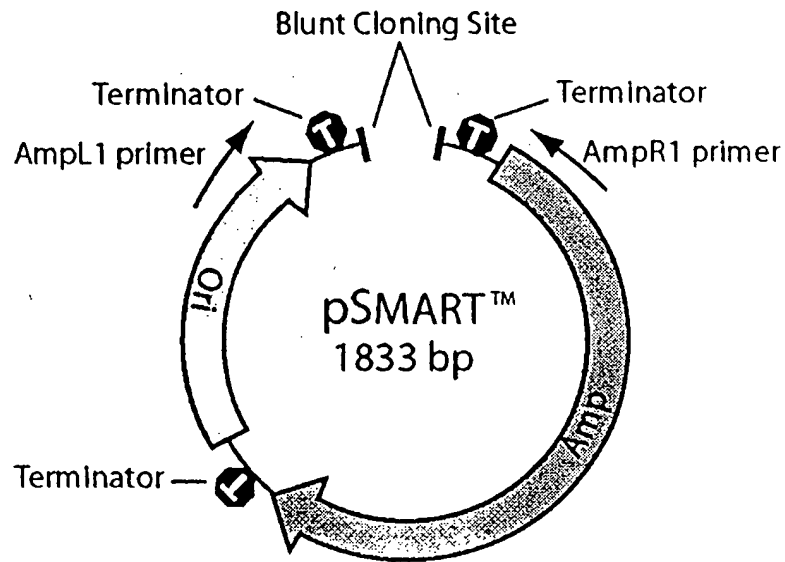


FIGURE 11 (SEQ ID NO:41)

TGATCGGCACGTAAGAGGTTCCAACCTTTCACCATAATGAAATAAGATCACTACCGGGCGTATTTTTTGGAGTTATCGAGAT
TTTCAGGAGCTAAGGAAGCTAAAATGGAGAAAAAAATCACTGGATATGCCACCGTTGATATATCCCAATGGCATCGTAAA
GAACATTTTGGAGGCATTTTCAGTCAGTTGCTCAATGTACCTATAACCAGACCGTTCAGCTGGATACTACGGCCTTTTAAA
GACCGTAAAGAAAAATAAGCACAAAGTTTTATCCGGCCTTTATTACATTCTTGCCCGCCTGATGAATGCTCATCCGGAAT
TCCGTATGGCAGTGAAAGACGGTGAGCTGGTGATATGGGATAGTGTTTACCCTTGTTACACCGTTTTCCATGAGCAAAC
GAAACGTTTTTCATCGCTCTGGAGTGAATACCACGACGATTTCCGGCAGTTTCTACACATATATTCGCAAGATGTGGCGTG
TTACGGTGAAAACCTGGCCTATTTCCCTAAAGGGTTTATTGAGAATATGTTTTTCGTCTCAGCCAATCCCTGGGTGAGTT
TCACCAGTTTTGATTTAAACGTGGCCAATATGGACAACTTCTTCGCCCCCGTTTTTACCATGGGCAATATTTATACGCAA
GGCGACAAGGTGCTGATGCCGCTGGCGATTTCAGGTTTCATCATGCCGTTTGTGATGGCTTCCATGTCGGCAGAATGCTTAA
TGAATTACAACAGTACTGCGATGAGTGGCAGGGCGGGCGTAAATGCAAAGCTTGTCATGCCCTGCAGGTCGACTCTAGAGG
ACCGGGTACCGAGCTCAGATCTTAGCATGGGACGTTTATATAGTGGTAATCTGGCAGCATTCAGGCAGCAACAAACAAG
CTGTTCCAGTTAGACTTAGCGGTCATTTATGATGACTGGTATAATGCCTATACAAGAAAAGATTGCATACGGTTACGTAT
TGAGGACAGATCTGGAAACCTGATTGATACTAGCACCTTCTACCACCACGACGAGGACGTTCTGTTCAATATGTGTACTG
ATTGGTTGAACCATATGTATGACCAGTTGAAGGACTGGAAGTAATAAGATCTTTGGTCCTGCAGGGAGCGTTAACATTTA
AATCGTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGA
AACCCGACAGGACTATAAAGATACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTCTCCTGTTCCGACCCCTGCCGCT
TACCGGATACCTGTCCGCCTTTCTCCCTTCGGGAAGCGTGGCGCTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCGG
TGTAGGTCGTTTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCCGTTTACGCCCCGACCGCTGCGCCTTATCCGGTAAC
TATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTA
TGTAAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGAACAGTATTTGGTATCTGCGCTC
TGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAAACAAACCACCGCTGGTAGCGGTGGTTTT
TTTGTGTTGCAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAGGATCGTTTGATCTTTTGGGCCC

FIGURE 12

A.



B.

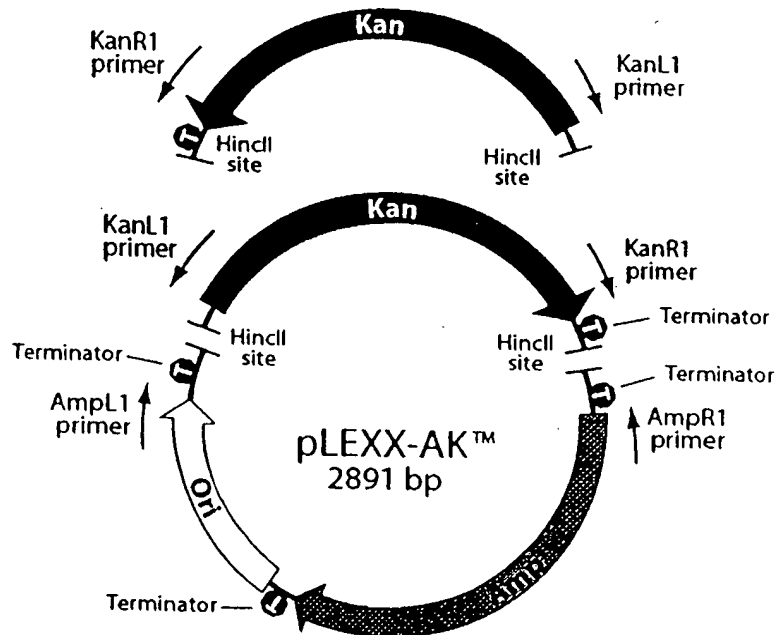


FIGURE 13

A. Kanamycin Gene Primers

(SEQ ID NO:114)
 KanL1 Primer → ATCTTGTCACGTCACATCAGAGATTTTGAGACACAAACGTTTAAACGGACGACGACCGGTATCGGCTGCAGGATCCAGATATGCGTC...
 TAGAACACGTTGCACTGTAGTCTCTAATAAATCTGTGTTGCAAAATTTGCCTGCTGCTGGCATAGCCGACGTCCTAGGTCTATACGCAG...
 Insert DNA

(SEQ ID NO:115)
 KanR1 Primer → TTCGTTTTTAATCTGGAAAAACCCACCCCTGGCGCTGCAGGTTCCAGATTCCCTGGTT...
 AAGCAAAATTAGACCTTTTGGTGGGACCGGACGTCCTCAAGGTCTAAGGACCAA...
 BstXI
 Insert DNA

B. Ampicillin Gene Primers

(SEQ ID NO:116)
 AmpL1 Primer → CAGTCCAGTTACGCTGGAGTCTGAGGCTCGTCTGTAATGATATCAAGCTTGAATTTCGTT...
 GTCAGGTCAATGCGACCTCAGACTCCGAGCAGGACTTACTATAGTTTCGAACTTAAGCAA...
 EcoRV HlnI III EcoRI
 Insert DNA

(SEQ ID NO:117)
 AmpR1 Primer → CTTTCTGCTATGGAGGTCAGGTATGATTTAAATGGTCAGTATTGACCGCATATCTAGAGAAATTCGTC...
 GAAAGACGATACCTCCAGTCCATATACTAAATTTACCAGTCATAACTCGCTATAGATCTCTTAAGCAG...
 EcoRV XbaI EcoRI
 Insert DNA

FIGURE 14

A. SEQ ID NO:85

GACGAATTCTCTAGATATCGCTCAATACTGACCATTTAAATCATACCTGA
CCTCCATAGCAGAAAGTCAAAAGCCTCCGACCGGAGGCTTTTGACTTGAT
CGGCACGTAAAGAGGTTCCAACCTTCCACCATAATGAAATAAGATCACTACC
GGGCGTATTTTTTGAGTTATCGAGATTTTCAGGAGCTAAGGAAGCTAAAA
TGAGTATTCAACATTTCCGTGTCGCCCTTATTCCTTTTTTGCGGCATTT
TGCCTTCCTGTTTTTGCTCACCAGAAACGCTGGTGAAAGTAAAGATGC
TGAAGATCAGTTGGGTGCACGAGTGGGTTACATCGAAGTGGATCTCAACA
GCGGTAAGATCCTTGAGAGTTTACGCCCCGAAGAACGTTTTCCAATGATG
AGCACTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGACGC
CGGGCAAGAGCAACTCGGTGCGCGCATACACTATTCTCAGAATGACTTGG
TTGAGTACTCACCAGTCACAGAAAAGCATCTCACGGATGGCATGACAGTA
AGAGAATTATGCAGTGCTGCCATAACCATGAGTGATAACACTGCGGCCAA
CTTACTTCTGGCAACGATCGGAGGACCGAAGGAGCTAACCGCTTTTTTGC
ACAACATGGGGGATCATGTAACCTCGCCTTGATCGTTGGGAACCGGAGCTG
AATGAAGCCATACCAAACGACGAGCGTGACACCACGATGCCTGTAGCAAT
GGCAACAACGTTGCGCAAACTATTAACCTGGCGAAGTACTTACTCTAGCTT
CCCGGCAACAATTAATAGACTGGATGGAGGCGGATAAAGTTGCAGGATCA
CTTCTGCGCTCGGCCCTCCCGGCTGGCTGGTTTTATTGCTGATAAATCTGG
AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATG
GTAAGCCCTCCCGCATCGTAGTTATCTACACGACGGGGAGTCAGGCAACT
ATGGATGAACGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAA
GCATTGTAATGAGGGCCAAATGTAATCACCTGGCTCACCTTCGGGTGG
GCCTTCTGCGTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAG
CATCACAAAAATCGATGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACT
ATAAAGATACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTCTCCTG
TTCCGACCCTGCGCTTACCGGATACCTGTCCGCCTTCTCCCTTCGGGA
AGCGTGCGCTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCCGGTGT
GGTCGTTGCTTCAAGCTGGGCTGTGTGCACGAACCCCCCGTTACGCCCCG
ACCGCTGCGCCTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGA
CACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGC
GAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACG
GCTACACTAGAAGAACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTT
ACCTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAAACAAACCCGCT
GGTAGCGGTGGTTTTTTTGTGTTGCAAGCAGCAGATTACGCGCAGAAAAA
AGGATCTCAAGAAGATCCTTTGATTTTCTACCGAAGAAAGGCCACCCGT
GAAGGTGAGCCAGTGAGTTGATTGCAGTCCAGTTACGCTGGAGTCTGAGG
CTCGTCTGAATGATATCAAGCTTGAATTCGTT

B. SEQ ID NO: 86

GACGCATATCTGGATCCTGCAGCCGATACGGTCGTCGTCCTTTAAACGTTGTGTCTCAAAATCTCTGATG
TCACGTTGCACAAGATAAAAAATATATCATCATGAACAATAAAACCGTCTGCTTACATAAACAGTAATACAA
GGGGTGTTATGAGCCATATTCAACGGGAACGCTTGTCTCGAGGCCGCGATTAAATTCCAACATGGATGCT
GATTTATATGGGTATAAATGGGCTCGCGATAATGTGGGCAATCAGGTGCGACAATCTATCGATTGTATGG
GAAGCCCGATGCGCCAGAGTTGTTTCTGAAACATGGCAAGGTAGCGTTGCCAATGATGTTACAGATGAGA
TGGTCAGGCTAAACTGGCTGACGGAATTTATGCCTCTTCCGACCATCAAGCATTTTATCCGTACTCCTGAT
GATGCATGGTTACTCACCCTGCGATCCAGGGAAAAACAGCATTCCAGGTATTAGAAGAATATCCTGATTC
AGGTGAAAAATATTGTTGATGCGCTGGCAGTGTCTCTGCGCCGGTTGCATTGATTCTCTGTTTGTAAATTGTC
CTTTTAAACGGCGATCGCGTATTTCTGCTCTGCTCAGGCGCAATCACGAATGAATAACGGTTTGGTTGGTGCG
AGTGATTTTGATGACGAGCGTAATGGCTGGCCTGTTGAACAAGTCTGGAAAGAAATGCATAAGCTTTTGCC
ATTCTACCCGGATTGATCGTCACTCATGGTGATTTCTCACTTGATAACCTTATTTTGGACGAGGGGAAAT
TAATAGGTTGATTGATGTTGGACGAGTCGGAATCGCAGACCGATACCAGGATCTTGCCATCCTATGGAAC
TGCCTCGGTGAGTTTTCTCCTTCAATTACAGAAACGGCTTTTCAAAAATATGGTATTGATAATCCTGATAT
GAATAAATGTCAGTTTCACTTGATGCTCGATGAGTTTTTCTAATCAAATAAAACGAAAGGCTCAGTCGAA
GACTGAGCCTTTCGTTTTAATCTGGA AAAACACCCTGGCGCTGCAGGTTCCAGATTCTCGGTT

Figure 15

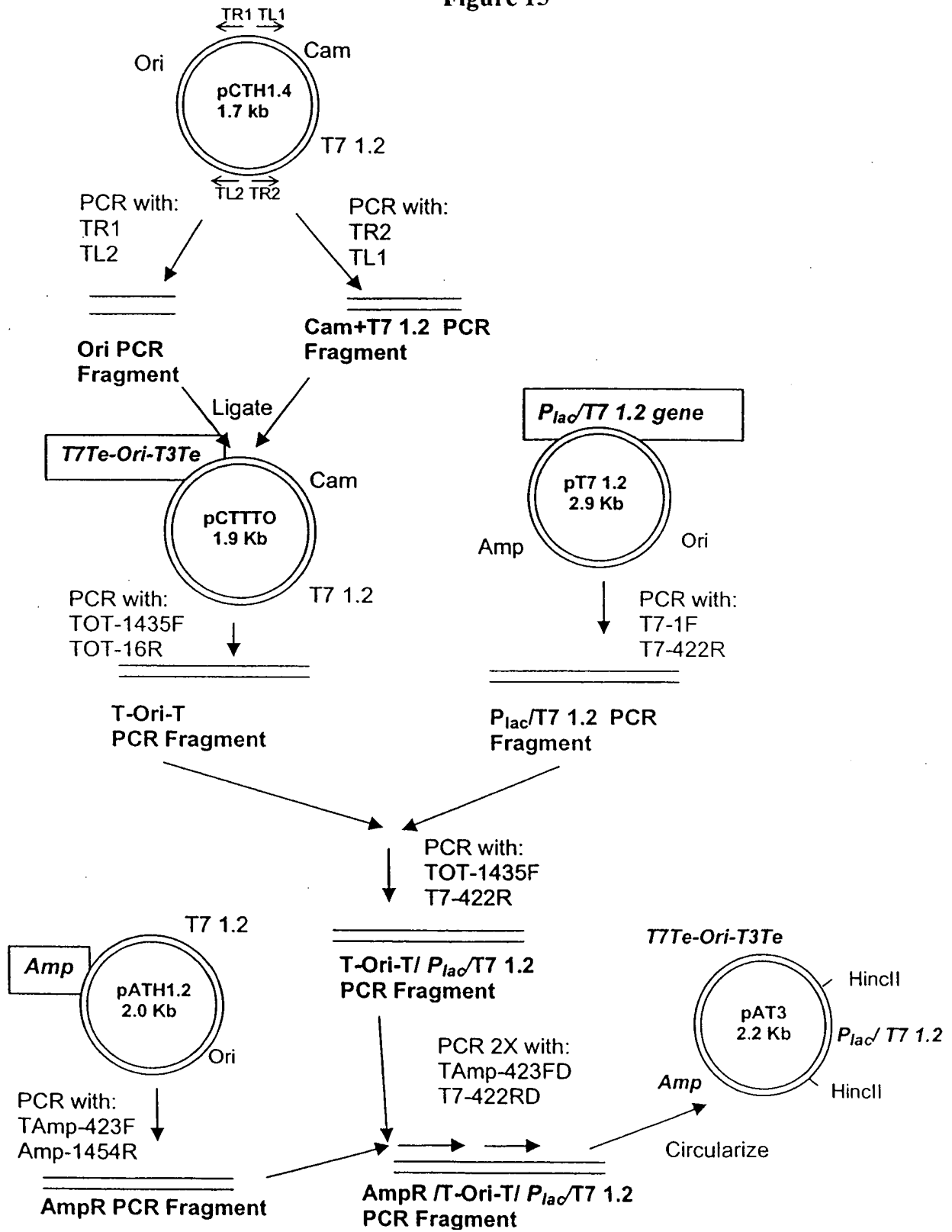


FIGURE 16

